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Reeb graph based examination of root development	064
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Abstract This paper presents an approach to analyze plant root development by means of topological image analysis. For phenotyping of plants their root development, the architecture of their root systems and thereby root characteristics such as branches and branch endings are analyzed. In order to simplify the examination of root characteristics and enable an efficient comparison of roots, a representation of imaged root data by Reeb graphs is introduced. Reeb graphs capture the topology of the represented structure in this case the locations of branches and branch endings of the roots - and form a skeletal representation of the underlying image data in this way. As the roots are pictured as 2D image data, the projection of a 3D structure to a 2D space might result in an overlap of branches in the image. One major advantage when analyzing roots based on Reeb graphs is posed by the ability to immediately distinguish between branching points and overlaps in the root structure. This is not as easily possible by an analysis solely based on contours.

## Introduction

Reeb graphs are widely used as shape descriptors for 3D structures. [2] gives a general overview on the use of Reeb graphs for shape analysis. [10] uses Reeb graphs for a pose independent segmentation of 3D data of human body scans, while [8] provides a skeletal representation of 2D data, Reeb graphs are for example used in [5] to provide a data skeletonization of the image content. However, Reeb graphs have not been applied to branched structures like roots or blood vessels although they pose a well suited representation. An analysis of branching patterns of roots based on a 3D reconstruction of the root architecture of rice plants is provided in [11].

<sup>-</sup> One of the ultimate challenges of biology is posed by the <sup>-</sup> question how genotypes translate into phenotypes. There,

082 the major bottleneck lies in the ability to phenotype a large .083 number of individuals and genotypes with high accuracy. This is particularly lagging in complex multicellular organ-.085 isms such as plants, in which specific biological processes often occur only temporarily and are restricted to specific .087 organs, tissues or even individual cells. Efficient and unsupervised image segmentation and the extraction of certain .089 characteristics are a key in approaching this goal. The .090 root of the small plant Arabidopsis thaliana is excellently .091 suitable for large-scale non-invasive phenotyping because \_092 it can be grown on transparent media in large numbers and \_093 its projections of the young root essentially capture all the 094 important biological features at the organ level. 095 When analyzing roots (for e.g. phenotyping), characteristics .096 such as the number of branches or the position and number .097 of branch-endings, are studied. These characteristics can be efficiently described by (Reeb) graphs. Reeb graphs .099 describe changes in topology in the represented structure. 100 Reeb graphs are based on Morse theory and analyze 101 the (here) image content according to a function (Morse 102 function). 103 When growing, roots change their shape, branches are 104 formed - their topology changes. Moreover the projection 105 of the 3D root structure to the 2D image data might cause .106 overlaps of branches in the image. In a Reeb graph a 107 distinction between a branch and an overlap is immediately 108 possible as these changes in topology are captured by the 109 graph. 110 The Reeb graph is used as a simplified, skeletal representa-.111 tion of the image data that captures the intrinsic topological 112 structure of the data and allows for a comparison of 113 the image content. Especially for the root dataset these 114 comparisons allow for a description of the growth process: 115

the image content. Especially for the root dataset these -113 comparisons allow for a description of the growth process: -114 the roots are imaged on consecutive days through their -115 growth period. In comparison with a simple standard -116 skeletonization approach as for example the Medial axis -117 transform, the skeleton derived by a Reeb graph not -118 only describes characteristics of the image content (here -120

121 branches of the roots) but captures the actual positions of 122 these characteristics as well. 123\_

124\_\_\_ The paper is structured as follows: Section 2 gives an 125 introduction to Reeb graphs, Section 3 describes the dataset used and Section 4 shows the computation of a Reeb graph 126 on the root dataset. The need for modifications of the Reeb 127 graphs and the types of modifications are discussed in 128 129\_ Section 5, Section 6 shows evaluation results on the root 130 dataset while a conclusion and a perspective to future work 131\_\_\_ are given in Section 7.

#### 133\_\_\_\_ 2 **Reeb Graphs and Morse Theory**

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134\_ Based on critical points according to a scalar function a 135 Reeb graph describes the topological structure that is the 136 connectivity of level sets of e.g. 2D or 3D content [4]. In 137. order to build a Reeb graph, critical points, of the structure 138. to be represented, need to be computed. 139

A point (a, b) of a function f(x, y) is called a critical point 140 if both derivatives  $f_x(a, b)$  and  $f_y(a, b)$  are equal 0 or if one 141. of these partial derivatives does not exist [9]. 142

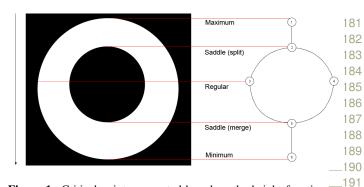
Such a critical point can either be a degenerate or-a 143\_ non-degenerate-critical point. These two cases can be 144 distinguished via the Hessian matrix. The determinant of 145\_ the Hessian matrix at a critical point x is then called the 146 discriminant. If this determinant is zero then x is called a 147 degenerate critical point of f (or non-Morse critical point of 148 f). Otherwise it is non-degenerate (or Morse critical point 149\_ of f).  $150_{-}$ 

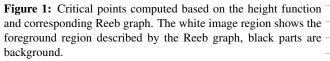
151\_ A smooth, real-valued function  $f: M_d \to \mathbb{R}$  is called a Morse function if it satisfies the following conditions for a d manifold  $M_d$  with or without boundary: 154\_

- all critical points of f are non-degenerate and lie inside  $M_d$ , 157\_\_\_\_
- 158\_ all critical points of f restricted to the boundary of  $M_d$ • 159\_ are non-degenerate
  - for all pairs of distinct critical points p and q,  $f(p) \neq p$ f(q) must hold [3].

Critical points of such a real-valued function are those points where the gradient becomes zero. The topological information of a shape described by a Reeb graph based on a function is related to the level sets of this function on the shape [2]. A change in topology appears with a change in the number of connected components in a level set. At regular points no topology changes occur. Topological changes occur at critical points only.

171. Reeb graphs are compact shape descriptors that preserve 172 the topological characteristics of the described shape [2]. 173 Vertices of the Reeb graph correspond to critical points of 174. the function (points where the topology of M changes), 175. edges describe topological persistence [2]. In other words: 176 All nodes having the same function value are represented by 177 one node in the graph, connections between nodes describe 178 connections between segments of the underlying structure. 179 Reeb graphs are originally defined for the continuous space, 180





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196 but have been extended to the discrete domain: Here the 197 Reeb graph is defined on a piecewise linear Morse function 198 [4]. As the approach presented in this paper provides an 199 analysis of 2D image content, it is based in the discrete 200 domain (image pixels). The Reeb graphs that are built 201 on the root images are therefore discrete Reeb graphs and 202 are based on the following definitions. In order to define 203 a discrete Reeb graph, connective point sets and level-set 204 curves are defined first: 205

- Two point sets are connected if there exists a pair of points (one point of each point sets) with a distance between these two points below a fixed threshold.
- If all non-empty subsets of a point set, as well as there complements, are connected, such a point set is called connective.
- A group of points that have the same Morse function value and that form a connective point set, is called a level-set curve.

The nodes in a discrete Reeb graph represent level-set curves, the edges connect two adjacent level-set curves, therefore the underlying point sets are connected [10].

222 In 2D critical points and corresponding nodes in the 223 Reeb graph are minima, maxima or saddles [3]. The 224 saddle nodes can be further distinguished: a saddle node that appears with a reduction in the number of connected 226 components is further called merge (saddle) node, a split 227 (saddle) node describes an increase in the number of connected components. When considering these two 229 different types of saddle nodes that might appear in a Reeb graph, four different types of critical points and according 231 nodes in the graph can be distinguished: maximum node, minimum node, split (saddle) node, merge (saddle) node. 233 Besides these nodes corresponding to critical points, regular 234 nodes can be added at any position and at-any edge in the Reeb graph as they do not describe a change in topology. 236 Nevertheless regular nodes can for example be used to 237 describe changes in the color of the foreground region (see 238 [1]). 239

241 The approach described in the following sections uses 242 the height function as Morse function  $\mu$ . In 2D the 243 height function is the function f that associates for each 244 point P = (x, y) the value y as the height of this point: 245  $f(x, y) \mapsto y$ .

Figure 1 shows an example for a Reeb graph based on a 246 height function, containing all five types of nodes and the 247 248\_\_\_ actual image the graph was computed on. Each edge in the 249\_\_\_ Reeb graph describes a connected component. Therefore 250 the edges of a Reeb graph are formed by connecting the 251\_\_\_ node representing the birth of a connected component 252\_\_\_ to the corresponding node representing the death of this 253\_\_\_ component.

## – 3 Root dataset

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257. For the root dataset images of the plant Arabidopsis thaliana 258. were taken. This plant is a model organism, which is widely 259 used in plant sciences, due to the small size of its genome, 260 the small size of the plant itself and its rapid life-cycle 261. [6]. The plants are grown on a nutrient containing agar gel 262 surface in plastic petri dishes that are vertically oriented. All 263 plants in one plate belong to one dataset. One dataset/plate 264 consists of 2 rows of 12 plants. The plates are placed in 265 a growth chamber that allows for controlled conditions as 266\_ constant temperature or humidity. 267.

The images are taken using an image scanner. A special 268. fixture allows for two datasets to be placed in an exact 269 known position inside the scanner. The images are acquired 270 with a scan at 1200 dpi resolution with 8bit color depth, 271 therefore one image is of approximately 6000x6000 pixels 272 in size. The images are stored as bmp files of about 150MB. 273 Along time several successive images are acquired this way, 274 as each plate is scanned at several successive days of the 275 growth process. A 3D stack of 2D images over time is thus 276. created for each root. 277\_

In a preprocessing step the 24 plants per plate are cropped to single images: one image per plant with an image size in the range of 500x1300 to 800x1300 pixels resolution and a file size of 1,5-2,5Mbyte. Example images of this dataset are shown in Figure 2.

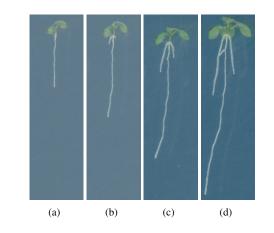
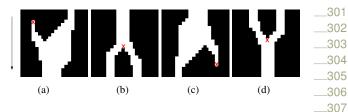


Figure 2: Example images of the root dataset: root004 - (a) day 8;
(b) day 12; (c) day 16; (d) day 20.



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**Figure 3:** Four different types of critical points computed according to the height function: (a) maximum / birth; (b) saddle (split); \_\_\_\_\_\_ (c) minimum / death; (d) saddle (merge).

The whole set of plant images used here consists of 9 sets of time series. Each set holds 6 images of one plant taken over time (day 1, day 4, day 8, day 12, day 16 and day 20 of the growth period). Of these 54 images, 34 images are analyzed, the other images are too early in the growth process and therefore to small in structure to be represented by a non-trivial Reeb graph.

All images analyzed are segmented in a preprocessing step – and consist of 2 foreground regions (leaves and roots, only – the roots are analyzed for this approach) and up to 2 holes – in the foreground structure. For reasons of the needed – preceded segmentation, the dataset is restricted in its size, – as the segmentation approach was done semi-automatically – and required a lot of time (up to 1.5h for one image).

### 4 Computation of Reeb graphs

As the roots are imaged in their natural direction of growth — (leaves in the top part of the image, roots growing downwards in a vertical direction) and branches occur mostly in this direction of growth, the height function is a suitable measuring function. Critical points indicate a change in topology, therefore they might only appear on the border of a region but not within the region. The borders of flat-regions in the image are analyzed to locate these critical points.

Figure 3 shows the four different types of critical points that are computed for the image content using a height function.

To compute the critical points a segmentation of the image needs to be done during a preprocessing step. As the height function is used to compute the critical points, the foreground region borders are analyzed with regard to horizontal borders as these might describe a change in the number of components. The so found critical points are located at the center of such a horizontal border.

There are two main problems encountered using this approach:

### 4.1 Critical points at same height

Due to the resolution of the image, the discretization of the root and further distortions during the segmentation process, it is possible that several critical points at different horizontal positions in the image are at the same vertical position (same height) in the image (see Figure 5(a) for an example). In this case the second criteria of Morse theory (see Section 360 361\_ 2) is not met. A Reeb graph cannot be built, as a decision on how to connect the nodes in order to build the graph can-362\_\_\_ 363\_\_\_ not be taken. Figure 4 shows an example: The solid lines 364 illustrate the only two fixed connections in this example, the 365 dashed lines indicate all possible connections. In this Reeb graph four edges are needed: one from each black (maxi-366 mum) node to a red (saddle) node and one edge from a red 367 368 (saddle) node to the green (minimum) node. A decision con-369\_\_\_ cerning these connections needs to be taken for the black 370 center node as well as for the two red nodes. A solution to build a Reeb graph, despite several critical points at the 371 372\_\_\_ same height, is discussed in Section 5.1

#### 373\_ 4.2 Additional critical points

374 Because of the segmentation prior to the computation of the 375 critical points, segmentation artefacts appear in the images. 376. The most common problem are frayed borders of image re-377. gions (see Figure 5(b) for an example). Especially for im-378. ages of day 16 the segmentation creates noise and distorted 379 region borders. When analyzing the images of day 16 one 380 notices a high humidity between the plates in the form of 381. water drops, which creates a highly texturized background 382 that complicates the segmentation. 383

These frayed borders in the segmented images result in addi-384 tional critical points that describe no actual split or merge of 385 the root structure. These artefacts alter the Reeb graph and 386. complicate a comparison or matching of graphs. One possi-387 bility on how to deal with these additional critical points is 388 described in Section 5.2. 389.

#### 5 Modifications on the graphs

To overcome the problems discussed in Section 4.1 and Section 4.2 the following techniques were used:

### 5.1 Controlled shift of critical point coordinates

Due to the discrete pixel-space, the coordinates x and y of a pixel p = (x, y) are integers. Critical points at the same height (same y-coordinate) occur for 35% of all images in the root dataset and are shifted. The height of such critical points is changed by an added factor  $f, 0 \leq f < 1$ . A critical point p = (x, y) is shifted to p' = (x, y + f), f is computed using the following formula:  $f = \frac{1}{w} \cdot (x-1)$ , with 403\_\_\_ w giving the width of the image. The y-coordinate is thereby 404\_\_\_

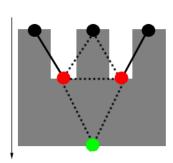


Figure 4: Critical points at same height: the solid lines show connections that are fixed, dashed lines indicate all possible connections - a decision needs to be taken for these.

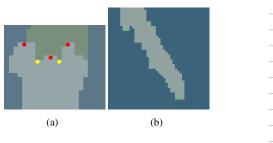


Figure 5: Problems encountered on the root dataset: (a) several critical points on same height; (b) frayed borders due to segmentation artifacts.

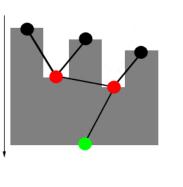
changed from an integer to a floating-point number. Critical points at the same height are moved downwards in a leftto-right order, thus for two critical points  $p_1 = (x_1, y)$  and  $p_2 = (x_2, y)$  with  $x_1 < x_2$ , it is valid that, after shifting the points to  $p'_1 = (x_1, y_1)$  and  $p'_2 = (x_2, y_2), y_1 < y_2$  holds. The actual order of heights is preserved by this correction procedure as only critical points that were primarily at the same height are changed. All critical points are at different heights, although when rounding down the y-coordinate of the critical points to an integer, they stay in the actual pixel line. A Reeb graph can therefore be built.

It is important to shift the heights in a fixed approach. A random decision choosing one of two critical points at the same height when building the Reeb graph cannot be used, as the results may vary with repeated tests. Reeb graphs built on such random decisions are not unique and therefore useless for e.g. comparison of two images.

Figure 6 shows a Reeb graph built on the marked critical points / nodes. Compared to Figure 4 where there are several critical points at the same height, Figure 6 shows critical points on different heights. The connections in this graph are unique. By shifting the critical points in Figure 4 according to the approach described in this section, the critical points are shifted to a configuration similar to the one shown in Figure 6.

## 5.2 Graph pruning

Due to the segmentation done as a preprocessing step, segmentation artefacts falsify the number of critical points and therefore the number of nodes and edges in the Reeb graph.



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Figure 6: Critical points at different heights, the connections in this Reeb graph are unique. \_\_\_480

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481	Number of nodes in graph					
482	Type of node	birth	split	merge	death	sum
483	no graph pruning	111	129	84	156	480
484	graph pruning	38	54	13	79	184
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**Table 1:** Total number of each type of nodes in the Reeb graphs of
 the root dataset with and without graph pruning.

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491\_\_\_ In order to use the extracted graphs as a skeletal representa-492 tion, branches that arise with artefacts need to be removed 493 from the Reeb graph.

494 For each pair of nodes adjacent nodes in the graph the Eu-495 clidean distance between these two nodes is computed. If 496 this distance is less than 1,5% of the image height such 497\_ connections are discarded and nodes are relinked if needed. 498\_ Regular nodes that may be introduced by this approach. As 499\_ these regular nodes do not contain any needed information, 500 they are removed after relinking. This threshold proved to 501\_ be the best choice in the experiments. This graph prun-502\_ ing results in a reduction of the overall number of nodes 503\_ in the Reeb graphs of the root dataset by 62%. Table 1 504\_ shows the numbers of nodes for all Reeb graphs in the root 505\_ dataset with and without graph pruning and Figure 7 shows 506\_ an example of the Reeb graph and the modified Reeb graph 507. for root 05, day 16. All the nodes in the lower part of the 508. root for the Reeb graph without graph pruning indicate spu-509 rious branches detected due to noise in the segmented im-510\_ age. These spurious branches are correctly discarded by the 511\_ graph pruning approach. 512\_

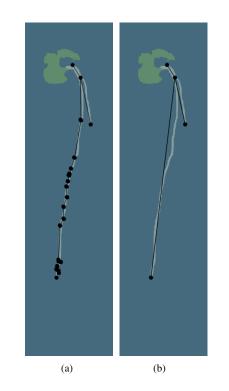


Figure 7: Reeb graph for root 05 day 16. (a) without graph pruning; (b) with graph pruning.

#### Results and evaluation on the root dataset 6

the Reeb graph.

overlap.

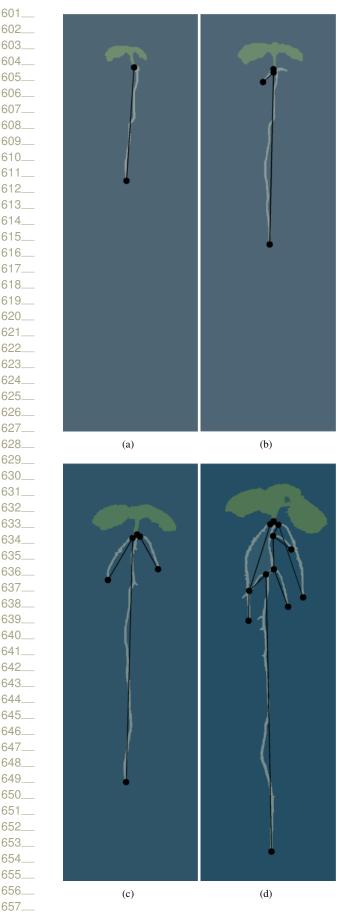
criteria have been evaluated:

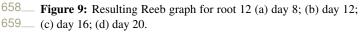
(a)

(c)

(c) day 16; (d) day 20.

541 542 543 Figure 8 shows the resulting Reeb graph for root 07 of 544 the dataset with both modifications implemented, drawn 545 as an overlay. There is a cycle in the Reeb graph for day 546 16 and day 20 (Figure 8(c) and 8(d)). In the image of day 547 12 (Figure 8(b)) there are three branches: The first and 548 the second branch overlap at sometime during the growth 549 process between day 12 and day 16. Because of this overlap .550 in the 3D space these two branches appear merged in the 551 2D projection of the image, therefore a cycle is formed in 552 553 554 Figure 9 shows the Reeb graphs for root 12 of the 555 dataset (both modifications are used). Some small branches 556 are not represented in the Reeb graphs of day 12, 16 and 20 557 as they resembled branches due to noise and were discarded 558 during the graph pruning process (see Section 6.1). Again a 559 cycle appears in the Reeb graph for day 20 as two branches 560 561 562 563 For the 34 single images of the root dataset the following .564 565 566 567 568 569 570 571 572 573 574 575 576 577 578 579 580 581 582 (b) 583 584 585 586 587 588 589 590 .591 592 593 594 595 596 (d) 597 598 Figure 8: Resulting Reeb graph for root 07 (a) day 8; (b) day 12; 599 600





wrong decisions on graph pruning			ing661
	images	false negatives	false positives662
graph pruning	8	10	0663
extension 1	18	4	36664
extension 2	13	10	9665
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**Table 2:** Branches wrongly discarded (false negative) and wrongly
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 accepted (false positive) in the graph pruning approach with two
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 different corrections based on pixel-color.
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# 6.1 Are all branches correctly detected and represented by the Reeb graph?

All major branches were correctly detected. Table 2 shows \_\_675 the number of images for which branches were wrongly dis- \_\_676 carded (false negatives) or wrongly accepted (false positive). \_\_677 For 23,5% of the images smaller branches were discarded \_\_678 due to the graph pruning as they resembled the frayed bor-\_\_679 680 small branches that describe actual root structures, their an-681 682 inscribe a larger angle than branches due to noise. However, -683 this assumption is based on the dataset presented and may -684 not be true for other datasets. Therefore another approach \_\_685 was tested: for a small branch with a critical point of type \_\_686 split, the color values at three pixels: at the critical point (a), — \_687 one row below the critical point (b) and two rows below the -\_688 critical point (c) were compared: 689 690

- 1. the color of (a) and (c) were taken from the segmented image, while the color value of (b) was taken from the unsegmented image —
- 2. all three color values were taken from the unsegmented \_\_\_696 image \_\_\_\_697

699 Branches are kept if the color value of (b) is closer to (a) than to (c). Table 2 shows the results for these two tests. 701 While the first option discards less true branches (false neg-702 atives) it keeps spurious branches for more than 50% of all 703 images. The second option keeps less spurious branches, 704 but does not reduce the number of false negatives compared to the graph pruning approach without these color compar-706 isons. Taking into account not only the color values of these 707 three pixels but of several neighbors, as it is done with Local 708 Binary Patterns, might present an option for future work. 709

# 6.2 Are additional branches (due to e.g. noise) detected?

As shown in Table 2 all additional branches (due to segmentation artefacts) are correctly discarded by the implemented \_\_\_\_713 graph pruning approach. \_\_\_\_715

For a series of images of one plant during the growth \_\_\_\_719 process the following factors have been analyzed: \_\_\_\_720

Number of nodes / edges / cycles				
	day 8	day 12	day 16	day 20
root 04	2/1/0	4/3/0	6/5/0	8/8/1
root 05	2/1/0	4/3/0	4/3/0	8/8/1
root 07	2/1/0	6/5/0	6/6/1	8/8/1
root 09	2/1/0	6/5/0	6/5/0	6/5/0
root 12	2/1/0	4/3/0	6/5/0	12/12/1
root 17	-	2/1/0	6/5/0	6/5/0
root 19	2/1/0	4/3/0	8/8/1	14 / 15 / 2
root 20	2/1/0	4/3/0	4/3/0	12/12/1
root 24	-	2/1/0	4/3/0	10/9/0

**Table 3:** Total number of nodes, edges and cycles in the modified graph (graph pruning without corrections) of each root image in the defined dataset.

## 6.3 Is an automatic grouping of images of one plant from different days possible?

As the roots grow downwards in a vertical direction, there are only minor changes in the position of the starting point of the actual root (transition between leaves an roots) - not accounting for actual movement of the plant (e.g. sliding down the plate). Therefore the starting point was used for this comparison. The average minimal Euclidean distance between all starting points is 14,4 pixels. Using this distance measurement to group one image of a root with earlier or later images of the same root, the grouping is correct for 71% of all images. However, images of day 16 falsify these numbers, as the plate of day 16 appears slightly enlarged in the image compared to the images of other days. As the images were automatically cut into single plant images in a preprocessing step, this scaling is not corrected. Excluding the images of day 16, the average minimal Euclidean distance decreases to 11,6 pixels and one image is grouped correctly with earlier or later images of the same root in 88%.

## - 6.4 General assumption: "Parts of a plant that appear in an early image of a plant do not disappear for a later image of the same plant."

This assumption proved to be correct for the images in the root dataset. The topology of a root only changes with the creation of new components (e.g. branches) over time. Table 3 shows the number of all nodes, edges and cycles in each (modified) graph of the root images.

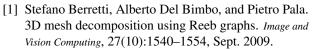
However, there is one exception to this assumption, which is based on the projection of a 3D structure to a 2D space. A branch in an early image of a plant might stay in the image of a later day, it may branch again but its ending may also disappear in the 2D image as it is merged with another branch due to an overlap of these two branches in the 3D space.

## 7 Conclusion and future work

Reeb graphs proved to be suitable descriptors for root structures as they capture the main characteristics of roots, namely branches and branch endings that are used in the phenotyping of plants, well. A Reeb graph provides a skeletal representation of a root that allows for fast analysis \_\_\_781 of root characteristics and efficient comparison of images \_\_782 and the contained root structure. Overlaps in 3D that appear \_\_\_783 as a merge of two branches in a 2D image are hard to \_\_784 distinguish from a branching point when analyzing only \_\_785 contours of image regions. Exploiting the topology of the \_\_786 root, actual branching points and overlaps in 3D can be \_\_787 immediately distinguished, as an overlap forms a cycle in \_\_788 the corresponding Reeb graph. 789 A future application in plant phenotyping is possible. \_ 790 However, for future work the segmentation approach \_\_791 needs to be changed to a less time-consuming (or even \_\_792 automatic) approach in order to allow for a larger dataset \_\_\_\_\_ to be analyzed. Moreover different functions will be used \_\_\_794 as Morse functions. Functions that should be taken into \_\_795 796 consideration are for example a medial axis as in [7] or \_\_\_\_ distance functions: for example the distance to a fixed point 797 in a structure, the sum of geodesic distance (both are used 798 in [10]) or the distance to an existing graph (as for example 799 a medial axis). 800 Open questions for future work (on the root dataset) are: 801 How does the chosen Morse function influence the correct 802 detection of branches in the root structure? Is the detection \_ 803 of all branches, respectively the detection of additional 804 branches due to noise, dependent on the Morse function \_\_805 used? Which Morse functions are able to correctly represent \_\_806 roots with a complex pattern of growth (e.g.: change in the \_\_807 main direction of growth)? 808 809

Furthermore this approach proofed to be suitable to extract \_ plant characteristics used in the phenotyping of plants, \_ \_

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